

OIPE

Raw Sequence Listing Error Summary

ERROR DETECTED SUGGESTED CORRECTION SERIAL NUMBER: 09/911,132

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1 Wrapped Nucleic
 Wrapped Aminos The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."
- 2 Invalid Line Length The rules require that a line not exceed 72 characters in length. This includes white spaces.
- 3 Misaligned Amino
 Numbering The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.
- 4 Non-ASCII The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.
- 5 Variable Length Sequence(s) _____ contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
- 6 PatentIn 2.0
 "bug" A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.
- 7 Skipped Sequences
(OLD RULES) Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence:
(2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
(i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
(xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
This sequence is intentionally skipped

Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.
- 8 Skipped Sequences
(NEW RULES) Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence.
<210> sequence id number
<400> sequence id number
000
- 9 Use of n's or Xaa's
(NEW RULES) Use of n's and/or Xaa's have been detected in the Sequence Listing.
Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
In <220> to <223> section, please explain location of n or Xaa; and which residue n or Xaa represents.
- 10 Invalid <213>
Response Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence
- 11 Use of <220>
 ✓ Sequence(s) ~~FFF~~ ^{most} missing the <220> "Feature" and associated numeric identifiers and responses.
Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.
(See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
- 12 PatentIn 2.0
 "bug" Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
- 13 Misuse of n n can only be used to represent a single nucleotide in a nucleic acid sequence. N is not used to represent any value not specifically a nucleotide.

0590
0906

#13



OIPE

ENTERED

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/911,132ADATE: 09/09/2002
TIME: 09:38:01Input Set : A:\RDID 0073US-2.ST25.txt
Output Set: N:\CRF4\09092002\I911132A.raw

3 <110> APPLICANT: Roche Diagnostics GmbH
 5 <120> TITLE OF INVENTION: Expression of Alkaline Phosphatase in Yeast
 7 <130> FILE REFERENCE: RDID 0073US
 9 <140> CURRENT APPLICATION NUMBER: 09/911,132A
 C--> 10 <141> CURRENT FILING DATE: 2002-08-28
 12 <160> NUMBER OF SEQ ID NOS: 38
 14 <170> SOFTWARE: PatentIn version 3.1
 16 <210> SEQ ID NO: 1
 17 <211> LENGTH: 1476
 18 <212> TYPE: DNA
 19 <213> ORGANISM: Bovine
 21 <400> SEQUENCE: 1

22	gaattcctca tcccagctga ggagggaaaac cccgccttct ggaaccgcca ggcagccag	60
24	gcccttgatg tagccaagaa gttcagccg atccagacag ctgccaagaa tgtcatcctc	120
26	ttcttggggg atggatggg ggtgcctacg gtgacagcaca ctcggatcct aaaggggcag	180
28	atgaatggca aactgggacc tgagacaccc ctggccatgg accagttccc atacgtggct	240
30	ctgtccaaga catacaacgt ggacagacag gtgccagacaa gcgcaggcac tgccactgcc	300
32	tacctgtgtg gggtaaaggg caactacaga accatcggtg taagtgcagc cgcccgctac	360
34	aatcagtgc acacgacacg tggaaatgag gtcacgtctg tgcataaccg ggccaagaaa	420
36	gcagggagg ccgtggaggt ggtgaccacc accagggtgc agcatgcctc cccagccggg	480
38	gcctacgcgc acacggtgaa ccgaaactgg tactcagacg ccgacctgcc tgctgatgca	540
40	cagaagaatg gctgccagga catgcgcgc cagctggct acaacatgga tattgacgtg	600
42	atcctgggtg gaggccaat gtacatgtt cctgagggga cccagaccc tgaataccca	660
44	gatgatgcca gtgtaatgg agtccggaa gacaaggcaga acctggtgca ggaatggcag	720
46	gccaaggcacc agggagccca gtatgtgtgg aaccgcactg cgctccttca ggcggccgat	780
48	gactccagtg taacacacacct catggccctc tttgagccgg cagacatgaa gtataatgtt	840
50	cagcaagacc acaccaagga cccgaccctg gcgagatga cggaggccgc cctgcaagtg	900
52	ctgagcagga accccccgggg cttctacctc ttctgtggagg gaggccgcat tgaccacggt	960
54	caccatgacg gcaaagctta tatggcactg actgaggcga tcatagttga caatgccatc	1020
56	gccaaggcta acgagctcac tagcgaactg gacacgctga tccttgtcac tgcagaccac	1080
58	tcccatgtct tctctttgg tggctacaca ctgcgtggga cctccatttt cggctctggcc	1140
60	cccgcaagg ctttagacag caagtcctac acctccatcc tctatggcaa tggcccaggc	1200
62	tatgcgttg gcgggggctc gagggccgat gttaatggca gcacaaggcga ggaaccctca	1260
64	taccggcagc aggccggccgt gcccctggct agcgagaccc acggggggcga agacgtggcg	1320
66	gtgttcgcgc gaggcccgca ggcgcacctg gtgcacggcg tgcaggagga gaccttcttg	1380
68	gcccacatca tggccttgc gggctgcgtg gagccctaca ccgactgcaa tctgccagcc	1440
70	cccgccaccg ccaccagcat cccgactag ggtacc	1476

73 <210> SEQ ID NO: 2
 74 <211> LENGTH: 40
 75 <212> TYPE: DNA
 76 <213> ORGANISM: Artificial Sequence
 78 <220> FEATURE:
 79 <223> OTHER INFORMATION: Primer

RAW SEQUENCE LISTING

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Input Set : A:\RDID 0073US-2.ST25.txt
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81 <400> SEQUENCE: 2
82 gcgcaattc ctcatccag ctgaggagga aaaccccgcc          40
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86 <211> LENGTH: 36
87 <212> TYPE: DNA
88 <213> ORGANISM: Artificial Sequence
90 <220> FEATURE:
91 <223> OTHER INFORMATION: Primer
93 <400> SEQUENCE: 3
94 cgcgggtacc ctatcgaaaa atgctggtgg cggtgg          36
97 <210> SEQ ID NO: 4
98 <211> LENGTH: 487
99 <212> TYPE: PRT
100 <213> ORGANISM: Bovine
102 <400> SEQUENCE: 4
104 Leu Ile Pro Ala Glu Glu Glu Asn Pro Ala Phe Trp Asn Arg Gln Ala
105 1           5           10           15
108 Ala Gln Ala Leu Asp Val Ala Lys Lys Leu Gln Pro Ile Gln Thr Ala
109           20           25           30
112 Ala Lys Asn Val Ile Leu Phe Leu Gly Asp Gly Met Gly Val Pro Thr
113           35           40           45
116 Val Thr Ala Thr Arg Ile Leu Lys Gly Gln Met Asn Gly Lys Leu Gly
117           50           55           60
120 Pro Glu Thr Pro Leu Ala Met Asp Gln Phe Pro Tyr Val Ala Leu Ser
121 65           70           75           80
124 Lys Thr Tyr Asn Val Asp Arg Gln Val Pro Asp Ser Ala Gly Thr Ala
125           85           90           95
128 Thr Ala Tyr Leu Cys Gly Val Lys Gly Asn Tyr Arg Thr Ile Gly Val
129           100          105          110
132 Ser Ala Ala Ala Arg Tyr Asn Gln Cys Asn Thr Arg Gly Asn Glu
133           115          120          125
136 Val Thr Ser Val Ile Asn Arg Ala Lys Lys Ala Gly Lys Ala Val Gly
137           130          135          140
140 Val Val Thr Thr Arg Val Gln His Ala Ser Pro Ala Gly Ala Tyr
141 145           150          155          160
144 Ala His Thr Val Asn Arg Asn Trp Tyr Ser Asp Ala Asp Leu Pro Ala
145           165          170          175
148 Asp Ala Gln Lys Asn Gly Cys Gln Asp Ile Ala Ala Gln Leu Val Tyr
149           180          185          190
152 Asn Met Asp Ile Asp Val Ile Leu Gly Gly Arg Met Tyr Met Phe
153           195          200          205
156 Pro Glu Gly Thr Pro Asp Pro Glu Tyr Pro Asp Asp Ala Ser Val Asn
157           210          215          220
160 Gly Val Arg Lys Asp Lys Gln Asn Leu Val Gln Glu Trp Gln Ala Lys
161 225           230          235          240
164 His Gln Gly Ala Gln Tyr Val Trp Asn Arg Thr Ala Leu Leu Gln Ala
165           245          250          255
168 Ala Asp Asp Ser Ser Val Thr His Leu Met Gly Leu Phe Glu Pro Ala
169           260          265          270

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172 Asp Met Lys Tyr Asn Val Gln Gln Asp His Thr Lys Asp Pro Thr Leu
 173 275 280 285
 176 Ala Glu Met Thr Glu Ala Ala Leu Gln Val Leu Ser Arg Asn Pro Arg
 177 290 295 300
 180 Gly Phe Tyr Leu Phe Val Glu Gly Gly Arg Ile Asp His Gly His His
 181 305 310 315 320
 184 Asp Gly Lys Ala Tyr Met Ala Leu Thr Glu Ala Ile Met Phe Asp Asn
 185 325 330 335
 188 Ala Ile Ala Lys Ala Asn Glu Leu Thr Ser Glu Leu Asp Thr Leu Ile
 189 340 345 350
 192 Leu Val Thr Ala Asp His Ser His Val Phe Ser Phe Gly Gly Tyr Thr
 193 355 360 365
 196 Leu Arg Gly Thr Ser Ile Phe Gly Leu Ala Pro Gly Lys Ala Leu Asp
 197 370 375 380
 200 Ser Lys Ser Tyr Thr Ser Ile Leu Tyr Gly Asn Gly Pro Gly Tyr Ala
 201 385 390 395 400
 204 Leu Gly Gly Ser Arg Pro Asp Val Asn Gly Ser Thr Ser Glu Glu
 205 405 410 415
 208 Pro Ser Tyr Arg Gln Gln Ala Ala Val Pro Leu Ala Ser Glu Thr His
 209 420 425 430
 212 Gly Gly Glu Asp Val Ala Val Phe Ala Arg Gly Pro Gln Ala His Leu
 213 435 440 445
 216 Val His Gly Val Gln Glu Glu Thr Phe Val Ala His Ile Met Ala Phe
 217 450 455 460
 220 Ala Gly Cys Val Glu Pro Tyr Thr Asp Cys Asn Leu Pro Ala Pro Ala
 221 465 470 475 480
 224 Thr Ala Thr Ser Ile Pro Asp
 225 485
 228 <210> SEQ ID NO: 5
 229 <211> LENGTH: 1476
 230 <212> TYPE: DNA
 231 <213> ORGANISM: Artificial Sequence
 233 <220> FEATURE:
 234 <223> OTHER INFORMATION: Codon-optimized DNA sequence
 236 <400> SEQUENCE: 5
 237 gaattcttga ttccagctga agaagaaaaat ccagctttt ggaatagaca agctgctcaa 60
 239 gctttggatg ttgctaagaa gttgcaacca attcaaactg ctgctaagaa ttttttttg 120
 241 tttttgggtg atggtatggg tttccaact gttactgcta ctagaattttt gaagggtcaa 180
 243 atgaatggta agttgggtcc agaaaactcca ttggctatgg atcaatttcc atacgttgct 240
 245 ttgtctaaga cttacaatgt tgatagacaa gttccagatt ctgctggtagt tgctactgct 300
 247 tacttgtgtg gtgttaaggg taattacaga actattggtg tttctgtgc tgctagatac 360
 249 aatcaatgta atactactag aggtaatgaa gttacttctg ttattaatag agctaagaag 420
 251 gctggtaagg ctgttgggtgt ttttactact actagagttc aacatgcttc tccagctgg 480
 253 gcttacgctc atactgttaa tagaaattgg tactctgatg ctgatttgcc agctgatgct 540
 255 caaaagaatg gttgtcaaga tattgctgct caattggttt acaatatgga tattgatgtt 600
 257 attttgggtg gtggtagaat gtacatgttt ccagaaggta ctccagatcc agaataccca 660
 259 gatgatgctt ctgttaatgg ttttagaaag gataagcaa atttgggtca agaatggcaa 720
 261 gcttaagcatc aaggtgctca atatgtttgg aatagaactg ctttgttgca agctgctgat 780
 263 gattcttagtg ttactcattt gatgggtttg tttgaaccag ctgatatgaa gtataatgtt 840

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Input Set : A:\RDID 0073US-2.ST25.txt
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265	caacaagatc atactaagga tccaaacttg gctgaaatga ctgaagctgc tttgcaagtt	900
267	ttgtctagaa atccaagagg ttttacttg tttgttgaag gtggtagaat tgatcatgg	960
269	catcatgatg gtaaggctta tatggcttg actgaagcta ttatgttga taatgctatt	1020
271	gctaaggcta atgaattgac ttctgaattg gatacttga tttgggtac tgctgatcat	1080
273	agtcatgtt tttctttgg tggttacact ttgagaggta cttctatTT tggTTggct	1140
275	ccaggttaagg ctttggatag taagtcttac acttctattt tgtatggtaa tggTccagg	1200
277	tatgcttgg gtggTgggtc tagaccagat gttaatggta gtactagtga agaaccatct	1260
279	tacagacaac aagctgctgt tccattggct agtgaardc atggTggta agatgttgct	1320
281	gttttgcta gaggtccaca agtcatttg gttcatggtg ttcaagaaga aactttgtt	1380
283	gctcatatta tggctttgc tggTTgttt gaaccataca ctgattgtaa tttgccagct	1440
285	ccagctactg ctactagtat tccagattaa ggtacc	1476
288	<210> SEQ ID NO: 6	
289	<211> LENGTH: 78	
290	<212> TYPE: DNA	
291	<213> ORGANISM: Artificial Sequence	
293	<220> FEATURE:	
294	<223> OTHER INFORMATION: Primer	
296	<400> SEQUENCE: 6	
297	gcgcgaattc ttgattccag ctgaagaaga aaatccagct ttttggata gacaagctgc	60
299	tcaagcttgc gatgttgc	78
302	<210> SEQ ID NO: 7	
303	<211> LENGTH: 70	
304	<212> TYPE: DNA	
305	<213> ORGANISM: Artificial Sequence	
307	<220> FEATURE:	
308	<223> OTHER INFORMATION: Primer	
310	<400> SEQUENCE: 7	
311	ccaaaaacaa aataacattc ttagcagcag tttgaattgg ttgcaacttc ttagcaacat	60
313	ccaaagcttgc	70
316	<210> SEQ ID NO: 8	
317	<211> LENGTH: 69	
318	<212> TYPE: DNA	
319	<213> ORGANISM: Artificial Sequence	
321	<220> FEATURE:	
322	<223> OTHER INFORMATION: Primer	
324	<400> SEQUENCE: 8	
325	gaatgttatt ttgttttgg gtgatggat ggggttcca actgttactg ctactagaat	60
327	tttgaaggg	69
330	<210> SEQ ID NO: 9	
331	<211> LENGTH: 70	
332	<212> TYPE: DNA	
333	<213> ORGANISM: Artificial Sequence	
335	<220> FEATURE:	
336	<223> OTHER INFORMATION: Primer	
338	<400> SEQUENCE: 9	
339	ggaaattgat ccatagccaa tggagttct ggacccaact taccattcat ttgacccttc	60
341	aaaattctag	70
344	<210> SEQ ID NO: 10	
345	<211> LENGTH: 71	

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346 <212> TYPE: DNA
347 <213> ORGANISM: Artificial Sequence
349 <220> FEATURE:
350 <223> OTHER INFORMATION: Primer
352 <400> SEQUENCE: 10
353 gctatggatc aattccata cggtgcttg tctaagactt acaatgtga tagacaagtt      60
355 ccagattctg c                               71
358 <210> SEQ ID NO: 11
359 <211> LENGTH: 71
360 <212> TYPE: DNA
361 <213> ORGANISM: Artificial Sequence
363 <220> FEATURE:
364 <223> OTHER INFORMATION: Primer
366 <400> SEQUENCE: 11
367 ccaatagttc tgtaattacc cttaacacca cacaagtaag cagtagcagt accagcagaa      60
369 tctggaaacctt g                           71
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373 <211> LENGTH: 72
374 <212> TYPE: DNA
375 <213> ORGANISM: Artificial Sequence
377 <220> FEATURE:
378 <223> OTHER INFORMATION: Primer
380 <400> SEQUENCE: 12
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383 gaggttatga ag                           72
386 <210> SEQ ID NO: 13
387 <211> LENGTH: 74
388 <212> TYPE: DNA
389 <213> ORGANISM: Artificial Sequence
391 <220> FEATURE:
392 <223> OTHER INFORMATION: Primer
394 <400> SEQUENCE: 13
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397 attacctcta gtag                           74
400 <210> SEQ ID NO: 14
401 <211> LENGTH: 74
402 <212> TYPE: DNA
403 <213> ORGANISM: Artificial Sequence
405 <220> FEATURE:
406 <223> OTHER INFORMATION: Primer
408 <400> SEQUENCE: 14
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411 catactgtta atag                           74
414 <210> SEQ ID NO: 15
415 <211> LENGTH: 68
416 <212> TYPE: DNA
417 <213> ORGANISM: Artificial Sequence
419 <220> FEATURE:
420 <223> OTHER INFORMATION: Primer

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VERIFICATION SUMMARY

PATENT APPLICATION: US/09/911,132A

DATE: 09/09/2002

TIME: 09:38:02

Input Set : A:\RDID 0073US-2.ST25.txt

Output Set: N:\CRF4\09092002\I911132A.raw

L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date